



## RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 09/553,431A  
Source: 1600  
Date Processed by STIC: 7/22/2003

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:  
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Revised 04/24/2003



1600

## RAW SEQUENCE LISTING

DATE: 07/22/2003

PATENT APPLICATION: US/09/553,431A

TIME: 13:26:19

Input Set : A:\Un900411.app

Output Set: N:\CRF4\07222003\I553431A.raw

3 <110> APPLICANT: Osteryoung, Katherine W.  
 5 <120> TITLE OF INVENTION: Manipulation of Min Genes in Plants  
 7 <130> FILE REFERENCE: 920905.90041  
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/553,431A  
 C--> 10 <141> CURRENT FILING DATE: 2000-04-19  
 12 <150> PRIOR APPLICATION NUMBER: 60/130,403  
 13 <151> PRIOR FILING DATE: 1999-04-19  
 15 <160> NUMBER OF SEQ ID NOS: 6  
 17 <170> SOFTWARE: PatentIn Ver. 2.1  
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 21 <212> TYPE: DNA  
 22 <213> ORGANISM: Arabidopsis thaliana  
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 26 <222> LOCATION: (1)..(978)  
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 33 cca tca tct ctc tca caa aag act cta ata tct tca cca aga ttc gtc 96  
 34 Pro Ser Ser Leu Ser Gln Lys Thr Leu Ile Ser Ser Pro Arg Phe Val  
 35 20 25 30  
 37 aat aac cct agc aga cgg agt cca ata cga tcc gtt ctt caa ttt aat 144  
 38 Asn Asn Pro Ser Arg Arg Ser Pro Ile Arg Ser Val Leu Gln Phe Asn  
 39 35 40 45  
 41 cgc aaa ccg gaa ctc gcc gga gaa acg ccg cgt atc gtc gtt atc acc 192  
 42 Arg Lys Pro Glu Leu Ala Gly Glu Thr Pro Arg Ile Val Val Ile Thr  
 43 50 55 60  
 45 tcc gga aaa ggc ggt gtt gga aag acg aca acc acc gca aat gtc ggt 240  
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 47 65 70 75 80  
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 51 85 90 95  
 53 ctt ggt ctc cgt aac ctc gat ctc ctc cta ggg tta gag aat cga gtc 336  
 54 Leu Gly Leu Arg Asn Leu Asp Leu Leu Gly Leu Glu Asn Arg Val  
 55 100 105 110  
 57 aat tac act tgc gtc gag gtt ata aac gga gat tgt cgt ctc gat caa 384  
 58 Asn Tyr Thr Cys Val Glu Val Ile Asn Gly Asp Cys Arg Leu Asp Gln  
 59 115 120 125  
 61 gct ctg gta cgt gat aag cgt tgg tcg aat ttc gaa ttg cta tgt ata 432  
 62 Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu Leu Cys Ile

Does Not Comply  
Corrected Diskette Needed

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Input Set : A:\Un900411.app

Output Set: N:\CRF4\07222003\I553431A.raw

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69 gaa tgg ctt gtg gat gcg ttg aaa act aga ccg gaa ggt tca ccg gat 528
70 Glu Trp Leu Val Asp Ala Leu Lys Thr Arg Pro Glu Gly Ser Pro Asp
71      165      170      175
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74 Phe Ile Ile Ile Asp Cys Pro Ala Gly Ile Asp Ala Gly Phe Ile Thr
75      180      185      190
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79      195      200      205
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82 Thr Ala Leu Arg Asp Ala Asp Arg Val Thr Gly Leu Leu Glu Cys Asp
83      210      215      220
85 gga atc aga gat ata aag atg att gtg aac aga gtg aga act gat atg 720
86 Gly Ile Arg Asp Ile Lys Met Ile Val Asn Arg Val Arg Thr Asp Met
87 225      230      235      240
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93 gga ttg tca ttg ctt ggt gta att cct gaa gat tct gag gtt att cga 816
94 Gly Leu Ser Leu Leu Gly Val Ile Pro Glu Asp Ser Glu Val Ile Arg
95      260      265      270
97 agc acg aat cga ggg ttt ccg ctt gtt ctg aat aag cct cct acg ctt 864
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99      275      280      285
101 gcg gga ttg gcg ttt gag cag gcg gct tgg aga ctc gtt gag caa gat 912
102 Ala Gly Leu Ala Phe Glu Gln Ala Ala Trp Arg Leu Val Glu Gln Asp
103      290      295      300
105 agt atg aag gct gtt atg gtg gag gaa gaa cct aag aaa cgt ggc ttc 960
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117 <213> ORGANISM: Arabidopsis thaliana
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124 20 25 30
126 Asn Asn Pro Ser Arg Arg Ser Pro Ile Arg Ser Val Leu Gln Phe Asn
127 35 40 45
129 Arg Lys Pro Glu Leu Ala Gly Glu Thr Pro Arg Ile Val Val Ile Thr

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Input Set : A:\Un900411.app

Output Set: N:\CRF4\07222003\I553431A.raw

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136      85      90      95
138 Leu Gly Leu Arg Asn Leu Asp Leu Leu Leu Gly Leu Glu Asn Arg Val
139      100      105      110
141 Asn Tyr Thr Cys Val Glu Val Ile Asn Gly Asp Cys Arg Leu Asp Gln
142      115      120      125
144 Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu Leu Cys Ile
145      130      135      140
147 Ser Lys Pro Arg Ser Lys Leu Pro Met Gly Phe Gly Gly Lys Ala Leu
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150 Glu Trp Leu Val Asp Ala Leu Lys Thr Arg Pro Glu Gly Ser Pro Asp
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154      180      185      190
156 Ala Ile Thr Pro Ala Asn Glu Ala Val Leu Val Thr Thr Pro Asp Ile
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159 Thr Ala Leu Arg Asp Ala Asp Arg Val Thr Gly Leu Leu Glu Cys Asp
160      210      215      220
162 Gly Ile Arg Asp Ile Lys Met Ile Val Asn Arg Val Arg Thr Asp Met
163 225      230      235      240
165 Ile Lys Gly Glu Asp Met Met Ser Val Leu Asp Val Gln Glu Met Leu
166      245      250      255
168 Gly Leu Ser Leu Leu Gly Val Ile Pro Glu Asp Ser Glu Val Ile Arg
169      260      265      270
171 Ser Thr Asn Arg Gly Phe Pro Leu Val Leu Asn Lys Pro Pro Thr Leu
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174 Ala Gly Leu Ala Phe Glu Gln Ala Ala Trp Arg Leu Val Glu Gln Asp
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188 <213> ORGANISM: Tagetes erecta
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Input Set : A:\Un900411.app

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207 tac cca aag cca cca cca att cgc tcc gtc ctt caa tac aat cgc aaa 202
208 Tyr Pro Lys Pro Pro Pro Ile Arg Ser Val Leu Gln Tyr Asn Arg Lys
209 40 45 50
211 cca gag ctc gcc gga gac act cca cga gtc gtc gca atc gac gcc gac 250
212 Pro Glu Leu Ala Gly Asp Thr Pro Arg Val Val Ala Ile Asp Ala Asp
213 55 60 65
215 gtt ggt cta cgt aac ctc gat ctt ctt ctc ggt ctc gaa aac cgc gtc 298
216 Val Gly Leu Arg Asn Leu Asp Leu Leu Gly Leu Glu Asn Arg Val
217 70 75 80
219 aat tac acc gtc gtt gaa gtt ctc aac ggc gat tgc aga ctc gac caa 346
220 Asn Tyr Thr Val Val Glu Val Leu Asn Gly Asp Cys Arg Leu Asp Gln
221 85 90 95
223 gcc cta gtt cgt gat aaa cgc tgg tca aat ttc gaa ttg ctt tgt att 394
224 Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu Leu Cys Ile
225 100 105 110 115
227 tca aaa cct agg tca aaa ttg cct tta gga ttt ggg gga aaa gct tta 442
228 Ser Lys Pro Arg Ser Lys Leu Pro Leu Gly Phe Gly Gly Lys Ala Leu
229 120 125 130
231 gtt tgg ctt gat gca tta aaa gat agg caa gaa ggt tgc ccg gat ttt 490
232 Val Trp Leu Asp Ala Leu Lys Asp Arg Gln Glu Gly Cys Pro Asp Phe
233 135 140 145
235 ata ctt ata gat tgt cct gca ggt att gat gcc ggg ttc ata acc gcc 538
236 Ile Leu Ile Asp Cys Pro Ala Gly Ile Asp Ala Gly Phe Ile Thr Ala
237 150 155 160
239 att aca ccg gct aac gaa gcc gta tta gtt aca aca cct gat att act 586
240 Ile Thr Pro Ala Asn Glu Ala Val Leu Val Thr Thr Pro Asp Ile Thr
241 165 170 175
243 gca ttg aga gat gca gat aga gtt aca ggc ttg ctt gaa tgt gat gga 634
244 Ala Leu Arg Asp Ala Asp Arg Val Thr Gly Leu Leu Glu Cys Asp Gly
245 180 185 190 195
247 att agg gat att aaa atg att gtg aac aga gtt aga act gat ttg ata 682
248 Ile Arg Asp Ile Lys Met Ile Val Asn Arg Val Arg Thr Asp Leu Ile
249 200 205 210
251 agg ggt gaa gat atg atg tca gtt ctt gat gtt caa gag atg ttg gga 730
252 Arg Gly Glu Asp Met Met Ser Val Leu Asp Val Gln Glu Met Leu Gly
253 215 220 225
255 ttg tca ttg ttg agt gat acc cga gga ttc gaa gtg att cgg agt acg 778
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257 230 235 240
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260 Asn Arg Gly Phe Pro Leu Val Leu Asn Lys Pro Pro Thr Leu Ala Gly
261 245 250 255
263 ttg gca ttt gag cag gct gct tgg aga ttg gtt gag caa gat agc atg 874
264 Leu Ala Phe Glu Gln Ala Ala Trp Arg Leu Val Glu Gln Asp Ser Met
265 260 265 270 275
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Output Set: N:\CRF4\07222003\I553431A.raw

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272 Phe Phe Gly Gly
273                295
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277 atgtgtacag taagctgata aggattgttt taattcagtt ttcagagaga aaattagaat 1094
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297          35          40          45
299 Asn Arg Lys Pro Glu Leu Ala Gly Asp Thr Pro Arg Val Val Ala Ile
300          50          55          60
302 Asp Ala Asp Val Gly Leu Arg Asn Leu Asp Leu Leu Gly Leu Glu
303   65          70          75          80
305 Asn Arg Val Asn Tyr Thr Val Val Glu Val Leu Asn Gly Asp Cys Arg
306          85          90          95
308 Leu Asp Gln Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu
309          100         105         110
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312          115         120         125
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327          195         200         205
329 Asp Leu Ile Arg Gly Glu Asp Met Met Ser Val Leu Asp Val Gln Glu
330          210         215         220
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333 225          230         235         240
335 Arg Ser Thr Asn Arg Gly Phe Pro Leu Val Leu Asn Lys Pro Pro Thr
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*see p. 7 for error explanation*

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/553,431A

DATE: 07/22/2003  
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Input Set : A:\Un900411.app  
Output Set: N:\CRF4\07222003\I553431A.raw

Use of <220> Feature (NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:5



**VERIFICATION SUMMARY**

DATE: 07/22/2003

PATENT APPLICATION: US/09/553,431A

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Input Set : A:\Un900411.app

Output Set: N:\CRF4\07222003\I553431A.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
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